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#### **RECOMBINANT PHAGE PROBES FOR *SALMONELLA TYPHIMURIUM* DETECTION**

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*Salmonella typhimurium* is a leading cause of inadvertent gastrointestinal foodborne illness in the United States. Although few actual accounts of deliberate food contamination have been documented in the United States, the recent advent of biocrimes and terrorism in our country suggests that this trend will not continue, highlighting the importance of rapidly identifying biological agents, regardless of the contamination origin, as one part of a comprehensive strategic plan to secure the public food supply. There is an urgent need for deployable, real-time threat agent detectors to replace traditional methods of food safety analysis that are slower, labor-intensive, and cost-inefficient. Confirmation of presence in food products can take as long as 48 hours by conventional culture. Current rapid detection initiatives include biosensors that routinely incorporate antibodies as the biorecognition unit. Although sensitive and specific, antibodies are costly and may degrade under unfavorable environmental conditions. We believe that a stable, inexpensive substitute for antibodies is filamentous phage manipulated through phage display technique then affinity selected for specificity to *S. typhimurium* from billion-clone phage landscape libraries. Our results show that recombinant phage affinity selected against *S. typhimurium* can be 12,000-22,000 times for more specific than controls and 10-1000 times more selective for *S. typhimurium* than

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other select enterobacteria. We anticipate that these highly specific, selective phage binders will build upon our current biosensor development initiatives for the rapid detection of biological agents such as *S. typhimurium*.

## Binding Confirmation

## Methods

## Phage specificity

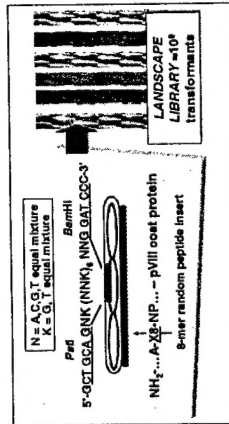
### Phage selectivity

**Salmonella typhimurium**

Amion, Acid Sensitive

Amion, Acid Sensitive

The diagram illustrates the phage selection protocol. It begins with a petri dish containing 'NZY agar' (labeled 'kan<sup>r</sup>') showing several phage plaques. An arrow points from this dish to another petri dish labeled 'E. coli (kan<sup>r</sup>)'. A large arrow then points to a box labeled 'Incubate 1 hour RT'. Another arrow points to a box labeled 'Wash 10X'. A final arrow points to a box labeled 'Lysate bound cells'.



### Phage selectivity

**Salmonella**  
specific  
phage E2

K2Y agar  
Antigen

*Salmonella enteritidis*  
*Salmonella typhimurium*

*Escherichia coli* (F<sup>-</sup>)  
*Proteus mirabilis*  
*Escherichia coli* serotypes  
*Citrobacter freundii*  
*Shigella flexneri*  
*Shigella sonnei*  
*Yersinia enterocolitica*  
*Yersinia pseudotuberculosis*  
*Serratia marcescens*  
*Pseudomonas aeruginosa*

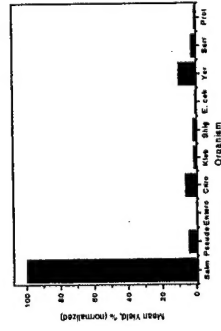
*E. coli* (kan<sup>r</sup>)

Figure 6 consists of two plots. The left plot shows the apparent rate constant  $k'$  (min<sup>-1</sup>) versus the A/UA ratio (mole/mole). The data points show a linear increase, and a linear fit is drawn through them. The right plot shows the apparent activation energy  $E_a$  (kJ/mole) versus the A/UA ratio (mole/mole). The data points show a non-linear decrease, and a curve is fitted to them.

A/UA (mole/mole)	$k'$ (min <sup>-1</sup> )	$E_a$ (kJ/mole)
0.0	0.0	10.0
0.2	0.2	9.5
0.4	0.4	9.0
0.6	0.6	8.5
0.8	0.8	8.0
1.0	1.0	7.5
1.2	1.2	7.0
1.4	1.4	6.5
1.6	1.6	6.0
1.8	1.8	5.5
2.0	2.0	5.0
2.2	2.2	4.5
2.4	2.4	4.0
2.6	2.6	3.5
2.8	2.8	3.0
3.0	3.0	2.5
3.2	3.2	2.0
3.4	3.4	1.5
3.6	3.6	1.0
3.8	3.8	0.5
4.0	4.0	0.0

**Average Phage Yield (CFU/ml)**

Determination of phage specificity and selectivity by binding of E2 to *S. typhimurium*. Determination of phage specificity at a concentration of  $10^9$  CFU/ml, resulted in low absolute yield percentage (= yield normalization). To determine if dose-dependent was a factor in conjunction with precipitation assay was performed by substituting the single concentrations ranging from  $10^{10}$  to  $10^7$  CFU/ml. A higher yield was demonstrated at  $10^8$  CFU/ml, when ratios of phage to cells approached 1.0, suggesting a bacteria interaction. Binding isotherm (Figure 5) confirms concentration-dependent interaction. The  $K_d$  apparent of the complex calculated from the Hill plot is  $3.18 \times 10^9$  estimated from Hill plot (Figure 6) is in good agreement with coefficient ( $n = 0.91$ )



**Fig. 2.** Precipitation assay demonstrated 90% greater affinity of phage **E2** (mean yield normalized) for *S. typhimurium* in comparison to challenge bacteria. Mean yield % is an average of 3 separate experiments normalized to a maximal mean yield of 2.8% from *S. typhimurium*.

## Acknowledgements

This work was supported by USDA grant No. 20013439410295C, and ARO/DARPA grant # DAAD 19-01-10454.

These results confirm our group's previous research efforts in the development of phage probes for the detection of biological molecules, commending landscape phage as substitute antibodies. We anticipate that these highly specific, selective phage binders will build upon our current biosensor development initiatives for the rapid detection of biological agents such as *S. typhimurium*.

## Conclusions

Product	Yield, %
CH <sub>4</sub>	0.05
C <sub>2</sub> H <sub>6</sub>	0.10
C <sub>2</sub> H <sub>4</sub>	0.15
C <sub>3</sub> H <sub>8</sub>	0.20
i-C <sub>4</sub> H <sub>10</sub>	0.25
n-C <sub>4</sub> H <sub>10</sub>	0.30
n-C <sub>5</sub> H <sub>12</sub>	0.35
n-C <sub>6</sub> H <sub>14</sub>	0.40
n-C <sub>7</sub> H <sub>16</sub>	0.45
n-C <sub>8</sub> H <sub>18</sub>	0.50
n-C <sub>9</sub> H <sub>20</sub>	0.55
n-C <sub>10</sub> H <sub>22</sub>	0.60
n-C <sub>11</sub> H <sub>24</sub>	0.65
n-C <sub>12</sub> H <sub>26</sub>	0.70
n-C <sub>13</sub> H <sub>28</sub>	0.75
n-C <sub>14</sub> H <sub>30</sub>	0.80
n-C <sub>15</sub> H <sub>32</sub>	0.85
n-C <sub>16</sub> H <sub>34</sub>	0.90
n-C <sub>17</sub> H <sub>36</sub>	0.95
n-C <sub>18</sub> H <sub>38</sub>	1.00
n-C <sub>19</sub> H <sub>40</sub>	1.05
n-C <sub>20</sub> H <sub>42</sub>	1.10
n-C <sub>21</sub> H <sub>44</sub>	1.15
n-C <sub>22</sub> H <sub>46</sub>	1.20
n-C <sub>23</sub> H <sub>48</sub>	1.25
n-C <sub>24</sub> H <sub>50</sub>	1.30
n-C <sub>25</sub> H <sub>52</sub>	1.35
n-C <sub>26</sub> H <sub>54</sub>	1.40
n-C <sub>27</sub> H <sub>56</sub>	1.45
n-C <sub>28</sub> H <sub>58</sub>	1.50
n-C <sub>29</sub> H <sub>60</sub>	1.55
n-C <sub>30</sub> H <sub>62</sub>	1.60
n-C <sub>31</sub> H <sub>64</sub>	1.65
n-C <sub>32</sub> H <sub>66</sub>	1.70
n-C <sub>33</sub> H <sub>68</sub>	1.75
n-C <sub>34</sub> H <sub>70</sub>	1.80
n-C <sub>35</sub> H <sub>72</sub>	1.85
n-C <sub>36</sub> H <sub>74</sub>	1.90
n-C <sub>37</sub> H <sub>76</sub>	1.95
n-C <sub>38</sub> H <sub>78</sub>	2.00
n-C <sub>39</sub> H <sub>80</sub>	2.05
n-C <sub>40</sub> H <sub>82</sub>	2.10
n-C <sub>41</sub> H <sub>84</sub>	2.15
n-C <sub>42</sub> H <sub>86</sub>	2.20
n-C <sub>43</sub> H <sub>88</sub>	2.25
n-C <sub>44</sub> H <sub>90</sub>	2.30
n-C <sub>45</sub> H <sub>92</sub>	2.35
n-C <sub>46</sub> H <sub>94</sub>	2.40
n-C <sub>47</sub> H <sub>96</sub>	2.45
n-C <sub>48</sub> H <sub>98</sub>	2.50
n-C <sub>49</sub> H <sub>100</sub>	2.55
n-C <sub>50</sub> H <sub>102</sub>	2.60
n-C <sub>51</sub> H <sub>104</sub>	2.65
n-C <sub>52</sub> H <sub>106</sub>	2.70
n-C <sub>53</sub> H <sub>108</sub>	2.75
n-C <sub>54</sub> H <sub>110</sub>	2.80
n-C <sub>55</sub> H <sub>112</sub>	2.85
n-C <sub>56</sub> H <sub>114</sub>	2.90
n-C <sub>57</sub> H <sub>116</sub>	2.95
n-C <sub>58</sub> H <sub>118</sub>	3.00
n-C <sub>59</sub> H <sub>120</sub>	3.05
n-C <sub>60</sub> H <sub>122</sub>	3.10
n-C <sub>61</sub> H <sub>124</sub>	3.15
n-C <sub>62</sub> H <sub>126</sub>	3.20
n-C <sub>63</sub> H <sub>128</sub>	3.25
n-C <sub>64</sub> H <sub>130</sub>	3.30
n-C <sub>65</sub> H <sub>132</sub>	3.35
n-C <sub>66</sub> H <sub>134</sub>	3.40
n-C <sub>67</sub> H <sub>136</sub>	3.45
n-C <sub>68</sub> H <sub>138</sub>	3.50
n-C <sub>69</sub> H <sub>140</sub>	3.55
n-C <sub>70</sub> H <sub>142</sub>	3.60
n-C <sub>71</sub> H <sub>144</sub>	3.65
n-C <sub>72</sub> H <sub>146</sub>	3.70
n-C <sub>73</sub> H <sub>148</sub>	3.75
n-C <sub>74</sub> H <sub>150</sub>	3.80
n-C <sub>75</sub> H <sub>152</sub>	3.85
n-C <sub>76</sub> H <sub>154</sub>	3.90
n-C <sub>77</sub> H <sub>156</sub>	3.95
n-C <sub>78</sub> H <sub>158</sub>	4.00
n-C <sub>79</sub> H <sub>160</sub>	4.05
n-C <sub>80</sub> H <sub>162</sub>	4.10
n-C <sub>81</sub> H <sub>164</sub>	4.15
n-C <sub>82</sub> H <sub>166</sub>	4.20
n-C <sub>83</sub> H <sub>168</sub>	4.25
n-C <sub>84</sub> H <sub>170</sub>	4.30
n-C <sub>85</sub> H <sub>172</sub>	4.35
n-C <sub>86</sub> H <sub>174</sub>	4.40
n-C <sub>87</sub> H <sub>176</sub>	4.45
n-C <sub>88</sub> H <sub>178</sub>	4.50</

**Fig. 1.** Specificity of select phage ( $10^6$  CFU/ml) were confirmed for *S. typhimurium* by phage capture, in comparison to ELISA (data not shown), which uses a solid support. Binding of best clones to *S. typhimurium* were 12,000–22,000 times greater than wild-type control phage 18–5, with phage clone E2 (VTP2PQHQ) possessing the highest binding efficiency among all select phage tested.

Since phase clone E2 demonstrated the greatest binding (% Yield) to *S. typhimurium*, we utilized this phage clone to visually confirm binding specificity. Fluorescent labeling of E2 using Alexa 488 fluorescent tag conferred an estimated 300 molecules of fluorochrome per phage particle. *S. typhimurium* incubated with the labeled phage was analyzed by fluorescence-activated cell sorting (FACS), TEM and fluorescence microscopy (data not shown).